

RAW SEQUENCE LISTING

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Application Serial Number: 09/986,625

Source: /FW16

Date Processed by STIC: 1/8/03

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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/986,625

DATE: 11/19/2004
TIME: 11:56:34

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Output Set: N:\CRF4\11192004\I986625.raw

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1 <110> APPLICANT: Hair, Gregory A.
2      Boden, Scott D.
3 <120> TITLE OF INVENTION: Novel Bone Mineralization Proteins, DNA, Vectors,
4      Expression Systems
5 <130> FILE REFERENCE: 06148.0115
6 <140> CURRENT APPLICATION NUMBER: 09/986,625
7 <141> CURRENT FILING DATE: 2001-11-09
8 <150> PRIOR APPLICATION NUMBER: US/09/124,238
9 <151> PRIOR FILING DATE: 1998-07-29
10 <150> PRIOR APPLICATION NUMBER: 60/054,219
11 <151> PRIOR FILING DATE: 1997-07-30
12 <150> PRIOR APPLICATION NUMBER: 60/080,407
13 <151> PRIOR FILING DATE: 1998-04-02
14 <160> NUMBER OF SEQ ID NOS: 35
15 <170> SOFTWARE: PatentIn Ver. 2.0
17 <210> SEQ ID NO: 1
18 <211> LENGTH: 457
19 <212> TYPE: PRT
20 <213> ORGANISM: Rattus norvegicus
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25      20         25          30
26 Leu Thr Pro Gly Gly Lys Ala Ala Gln Ala Gly Val Ala Val Gly Asp
27      35         40          45
28 Trp Val Leu Ser Ile Asp Gly Glu Asn Ala Gly Ser Leu Thr His Ile
29      50         55          60
30 Glu Ala Gln Asn Lys Ile Arg Ala Cys Gly Glu Arg Leu Ser Leu Gly
31      65         70          75          80
32 Leu Ser Arg Ala Gln Pro Ala Gln Ser Lys Pro Gln Lys Ala Leu Thr
33      85         90          95
34 Pro Pro Ala Asp Pro Pro Arg Tyr Thr Phe Ala Pro Ser Ala Ser Leu
35      100        105         110
36 Asn Lys Thr Ala Arg Pro Phe Gly Ala Pro Pro Pro Thr Asp Ser Ala
37      115        120         125
38 Leu Ser Gln Asn Gly Gln Leu Leu Arg Gln Leu Val Pro Asp Ala Ser
39      130        135         140
40 Lys Gln Arg Leu Met Glu Asn Thr Glu Asp Trp Arg Pro Arg Pro Gly
41      145        150         155         160
42 Thr Gly Gln Ser Arg Ser Phe Arg Ile Leu Ala His Leu Thr Gly Thr
43      165        170         175
44 Glu Phe Met Gln Asp Pro Asp Glu Glu Phe Met Lys Lys Ser Ser Gln

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46 Val Pro Arg Thr Glu Ala Pro Ala Pro Ala Ser Thr Ile Pro Gln Glu
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48 Ser Trp Pro Gly Pro Thr Thr Pro Ser Pro Thr Ser Arg Pro Pro Trp
49           210           215           220
50 Ala Val Asp Pro Ala Phe Ala Glu Arg Tyr Ala Pro Asp Lys Thr Ser
51           225           230           235           240
52 Thr Val Leu Thr Arg His Ser Gln Pro Ala Thr Pro Thr Pro Leu Gln
53           245           250           255
54 Asn Arg Thr Ser Ile Val Gln Ala Ala Ala Gly Gly Gly Thr Gly Gly
55           260           265           270
56 Gly Ser Asn Asn Gly Lys Thr Pro Val Cys His Gln Cys His Lys Ile
57           275           280           285
58 Ile Arg Gly Arg Tyr Leu Val Ala Leu Gly His Ala Tyr His Pro Glu
59           290           295           300
60 Glu Phe Val Cys Ser Gln Cys Gly Lys Val Leu Glu Glu Gly Gly Phe
61           305           310           315           320
62 Phe Glu Glu Lys Gly Ala Ile Phe Cys Pro Ser Cys Tyr Asp Val Arg
63           325           330           335
64 Tyr Ala Pro Ser Cys Ala Lys Cys Lys Lys Ile Thr Gly Glu Ile
65           340           345           350
66 Met His Ala Leu Lys Met Thr Trp His Val Pro Cys Phe Thr Cys Ala
67           355           360           365
68 Ala Cys Lys Thr Pro Ile Arg Asn Arg Ala Phe Tyr Met Glu Glu Gly
69           370           375           380
70 Ala Pro Tyr Cys Glu Arg Asp Tyr Glu Lys Met Phe Gly Thr Lys Cys
71           385           390           395           400
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73           405           410           415
74 Leu Gly Phe Ser Trp His Asp Thr Cys Phe Val Cys Ala Ile Cys Gln
75           420           425           430
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81 <210> SEQ ID NO: 2

82 <211> LENGTH: 1696

83 <212> TYPE: DNA

84 <213> ORGANISM: Rattus norvegicus

85 <400> SEQUENCE: 2

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88 cgtctgcaag gggcaagga cttcaacgtg cccctctcca tctctcggtt cactcctggaa 180
89 ggcaaggccg cacaggccgg tgtggccgtg ggagactggg tactgagtat cgacggtgag 240
90 aacgcccggaa gcctcacaca cattgaagcc cagaacaaga tccgtgcctg tggggagcgc 300
91 ctcagcctgg gtcttagcag agcccagcct gctcagagca aaccacagaa ggcctgacc 360
92 cctcccgccg accccccgag gtacacttt gcaccaagcg cctccctcaa caagacggcc 420
93 cggcccttcg gggcacccccc acctactgac agccctgtt cgcagaatgg acagctgctc 480
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97  gaagccccag ccccaagcctc aaccataccc caggaatcc ggcctggccc caccaccccc 720
98  agccccacca gcccggccacc ctggggcgta gatcctgcat ttgctgagcg ctatgcccc 780
99  gacaaaacca gcacagtgt gacccgacac agccagccag ccacacctac gcctctgcag 840
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102 ctggggccacg cgtaccatcc tgaggaattt gtgtgcagcc agtgtggaa ggtcctggaa 1020
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112 cctggccaga tcctggggct ccctcctcac agtccccctt cccacactc ctccaccacc 1620
113 accaccgtca ctcacagggtg ctgcctctt agccccagtt cactctgggtg tcacaataaa 1680
114 cctgtatgtt gctgtt 1696

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123 ttcagctggc atgatacggt ttttgcgaa atatcaactt ggaaggaaag 180
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127 <210> SEQ ID NO: 4
128 <211> LENGTH: 16
129 <212> TYPE: DNA
130 <213> ORGANISM: Artificial Sequence
131 <220> FEATURE:
132 <223> OTHER INFORMATION: Differential Display PCR Primer
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138 <212> TYPE: DNA
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141 <223> OTHER INFORMATION: Differential Display PCR Primer
142 <400> SEQUENCE: 5
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146 <211> LENGTH: 223
147 <212> TYPE: DNA

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 156 <211> LENGTH: 717
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 167 ccagatgcca gcaagcagcg gctgatggag aacacagagg actggcggcc gggccgggg 480
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 193 atgcacgcgc tgaagatgac ctggcacgtc cactgcttta cctgtgtgc ctgcaagacg 960
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 247 Leu Thr Pro Gly Gly Lys Ala Ala Gln Ala Gly Val Ala Val Gly Asp
 248 35 40 45
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